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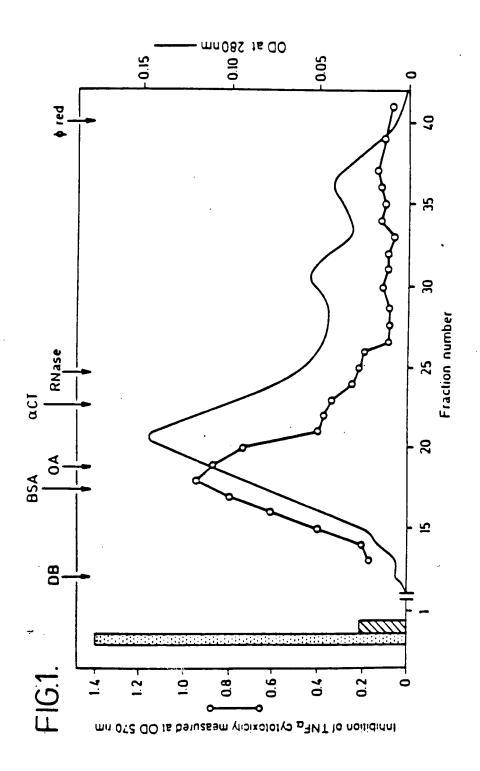
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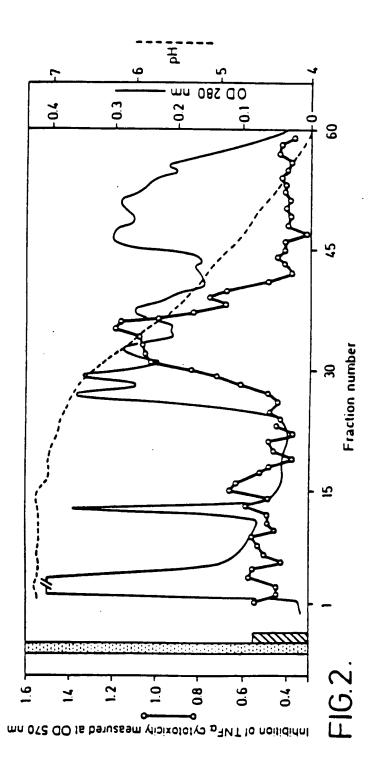
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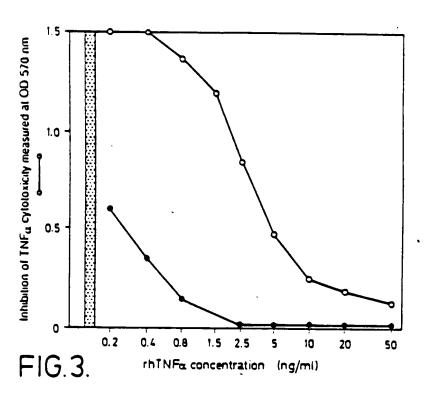
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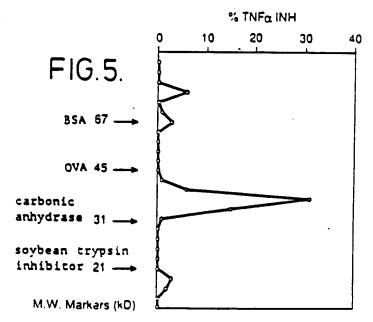
(54) TNF-elpha inhibitors

(57) A protein having a selective tumour necrosis factor (TNF)-α inhibitory activity, but which does not block other proteins. particularly IL-1, is prepared by extraction from urine of febule patients.

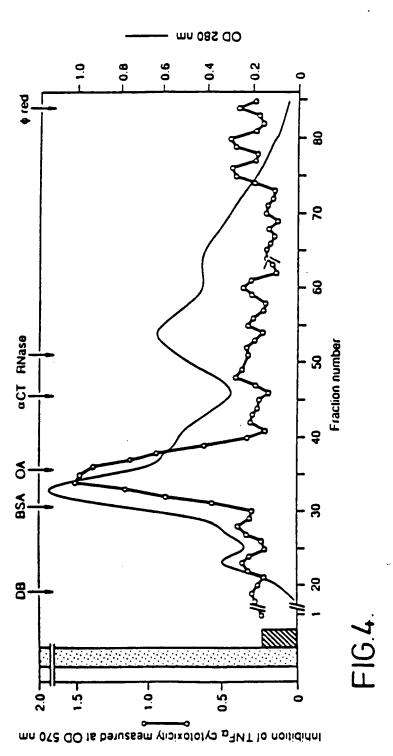


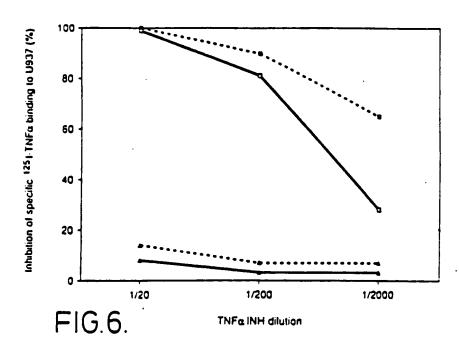


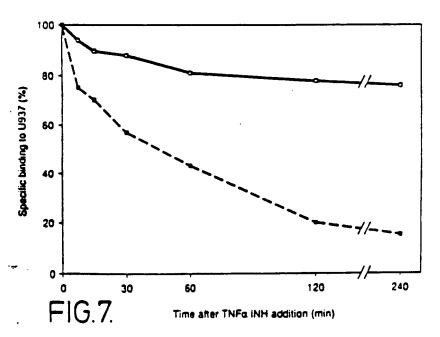












BIOLOGICALLY ACTIVE PROTEINS

The present invention relates to a novel protein having an inhibitory effect against Tumour Necrosis Factor α -mediated activity, to the isolation and purification of such a protein from natural sources, to its preparation by DNA manipulation and to the use of such a protein in the treatment of conditions associated with excessive or unregulated TNF α production.

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Tumour necrosis factor (TNF) is an activity embodied by a family of at least two proteins, α and β, which are cytotoxic for tumour cells and inhibit their growth in culture [E. Carswell et. al. "An endotoxin-induced serum factor that causes necrosis of tumours", Proc. Natl. Acad. Sci. USA, 72, p3666 (1975)]. Tumour necrosis factor α (TNFα), also termed "cachectin", is mainly produced by cells of the monocyte/macrophage lineage in response to "stress" signals which accompany invasive stimuli such as bacteria, viruses, tumours and other toxins. TNFβ, commonly termed "lymphotoxin", is mainly croduced by lymphoid cells. TNFβ has many activities similar to those of TNFα, but it appears to be less potent although this may be as a result of difficulties in preparing pure TNFβ.

activities [8. Beutler et. al., "Identity of tumour necrosis factor and the macrophage-secreted factor cachectin", Nature, 316, p552 (1985)] sharing several of them with interleukin 1 (IL-1) [J. Le et. al., "Tumour necrosis factor and interleukin 1: cytokines with multiple overlapping biological activities", Laboratory Invest., 56, p234 (1987)]. Elevated levels of INFa induced by, for example, tumour

cells may lead to weight loss and cachexia and TNFα has also been implicated as a principal mediator of endotoxic shock (septic shock) which can be fatal. Other biological effects of TNFα include hypotension, fever (induced by stimulation of hypothalamic prostoglandin E₂ (PGE₂) synthesis), coagulopathy (induced by stimulation of vascular endothelial cells which release, for example, tissue factor) and tissue destruction (induced by, for example, stimulation of a series of proteinases, including collagenase production by dermal fibroblasts and synovial cells) [C. Dinarello et. al., "Tumour necrosis factor (cachectin) is an endogenous pyrogen and induces production of interleukin 1", J. Exp. Med., 163, p1433 (1986); J. Dayer et. al., "Cachectin/tumour necrosis factor stimulate collagenase and prostaglandin E₂ production by human dermal fibroblasts and synovial cells", J. Exp. Med., 162, p2163 (1985)].

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There exists, therefore, a need to develop a cachectin/TNFz inhibitor which prevents endotoxic shock, cachexia and the other deleterious effects described above. It has been shown that passive immunisation of animals against cachectin can prevent endotoxin-induced death, mediated by TNF α antibodies [8. Beutler et. al., Nature, 316, supra].

We have now identified a novel protein which has a potent inhibitory effect against $TNF\alpha$ -mediated activities without significant concomitant inhibition of IL-1-mediated activity. The protein is hereinafter identified as Tumour Necrosis Factor α Inhibitor ($TNF\alpha$ INH).

. Thus in one aspect of the invention, we provide a protein which selectively inhibits tumour necrosis factor α -mediated activity.

As used herein, selective inhibition as shown by the innibitor of the invention is identified as the ability to block TNF-mediated activity while lacking the ability to block other proteins which have in common with TNF certain but not all of the biological activities of TNF, such as IL-1.

Preferably, the tumour necrosis factor a inhibitor of the invention is in a substantially homogeneous form, being substantially free from major contaminents and/or substantially free from other proteinaceous material.

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The tumour necrosis factor α inhibitor according to the invention has been found to have one or more of the following characteristics:

- (a) a molecular weight in the range 40 to 60 kDa, determined by molecular sieve chromatography;
- (b) an iso-electric point (pI) in the range 5.5 to 6.1, determined by chromatofocussing;
- (c) inhibition of the standard TNF assay of differential cytotoxicity for murine L929 cells treated with actinomycin D, as described by G. Nedwin et. al. "Effects of interleukin 2, interferon-γ and mitogens on the production of tumour necrosis factors α and ε", J. Immunol., 135, p2492 (1985). This inhibition can be overcome by further addition of TNFα, indicating that the inhibition is competitive. The inhibitor is also an inhibitor of TNFβ activity, although inhibition of TNFα in this assay is more efficient than that of JNFβ;
- (d) inhibition of TNF-induced PGE $_2$ release from human fibroblasts and synovial cells;

- (e) the inhibitor interferes with the binding of TNF2 to UPP7 cells (a monocytic tumour line) as evidenced by inhibition of binding of radiolabelled TNF α (12 I-TNF α);
- (f) the dissociation of a pre-formed TNFa : U937 cell complex is promoted by the inhibitor in a temperature dependent manner;
 - (g) the inhibitor does not degrade TNF by proteolytic cleavage;

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(h) the inhibitor does not inhibit IL-1 receptor-binding activity e.g. the binding of radiolabelled IL-1 (12 I-IL-1 α) to the murine thymoma subline EL4-6.1.

We have found that the protein of the invention when further purified has a molecular weight of about 33000 daltons as determined by sodium dodecyl sulphate polyacrylamide gel electrophoresis (SDS PAGE).

- There is thus provided as a further or alternative aspect, a protein which selectively inhibits $INF\alpha$ -mediated activity which has one or more of the following characteristics:
 - (a) a molecular weight of about 33 kDa, determined by SDS PAGE:
 - (b) an iso-electric point (pI) in the range 5.5 to 6.1, determined by chromatofocussing;
 - (c) inhibition of the standard TNF assay of differential cytotoxicity for murine L929 cells treated with actinomycin D, as described by G. Nedwin et. al. "Effects of interleukin 2, interferon-y and mitogens on the production of tumour necrosis factors α and β ", β . Immunol., 135, p2492 (1985). This inhibition can be overcome by further addition of TNF α , indicating that the inhibition is

competitive. The inhibitor is also an inhibitor of INFS activity, although inhibition of INF α in this assay is more efficient than that of INFS;

- (d) inhibition of TNF-induced PGE release from human fibroblasts and synovial cells:
 - (e) the inhibitor interferes with the binding of INF_x to U937 cells (a monocytic tumour line) as evidenced by inhibition of binding of radiolabelled INF_x ($^{1/2}$ I-INF $_x$);
- (f) the dissociation of a pre-formed TNF α : U937 cell complex is promoted by the inhibitor in a temperature dependent manner:
 - (g) the inhibitor does not degrade TNF by proteolytic cleavage;
- (h) the inhibitor does not inhibit IL-1 receptor-binding activity e.g. the binding of radiolabelled IL-1 (12 I-IL-1 $_{\alpha}$) to the murine thymoma subline EL4-6.1.

Preferably the TNF α INH of the present invention has both of the characteristics (a) and (b) and one or more of the characteristics (c) to (h).

In particular, the TNF α INH of the present invention has all of the characteristics (a) to (h).

The protein of the invention has an amino terminal amino acid sequence as follows:

Asp-Ser-Val-Cys-Pro-Gln-Gly-Lys-Tyr-Ile-His-Pro-Gln-Cys-Asn-Ser-Ile

It is further believed that the next three amino acids provide a glycosylation site and that the sequence thus continues is in-Ser-Thr-Lys.

It will be appreciated that a INFx inhibitor according to the invention will comprise an amino acid sequence substantially corresponding to the sequence of native INFx INH and containing an amino terminal sequence substantially identical to that described above. The sequence of a INFx inhibitor according to the invention will thus be identical to the sequence of native INFx INH or contain one or more deletions, substitutions, insertions, inversions or additions of allelic origin or otherwise, the resulting sequence will have at least 80% and preferably 90%, homology with the sequence of native INFx INH and retain essentially the same biological properties of the protein.

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The TNFs inhibitor of the invention has been demonstrated to be proteinaceous in that it is inactivated by heating in a time and temperature dependent manner is is destroyed by treatment with trypsin or promase.

The INF α INH of the invention has also been shown to be a glycoprotein since treatment with the enzyme Endoglycosidase F reduces the molecular weight by 7 to 8 kDa.

In a further or alternative aspect of the invention there is thus provided a $INF\alpha$ inhibitor as defined herein, but which is in a substantially unglycosylated state.

The inhibitors of the invention are of interest in the treatment of conditions in which it is desirable to inhibit $TNF\alpha$ activity, for example, those which arise from the effects of $TNF\alpha$ such as weight loss, shock, cachexia and chronic local inflammation, rheumatoid arthritis, disseminated intravascular coagulation and nephritis.

There is thus provided as a further aspect of the invention a TNF2 inhibitor as berein defined or a pharmaceutically acceptable derivative thereof for use as an active therapeutic agent, in particular, in the treatment of conditions associated with excessive or unregulated TNF2 production.

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In a further or alternative aspect of the invention there is provided a method for the treatment of conditions associated with excessive or unregulated TNF α production in a mammal including man comprising administration of an effective amount of a TNF α inhibitor as herein defined or a pharmaceutically acceptable derivative thereof.

There is also provided in a further or alternative aspect of the invention use of a TNFz inhibitor as herein defined or a pharmaceutically acceptable derivative thereof for the manufacture of a medicament for the treatment of conditions associated with excessive or unregulated TNFz production.

It will be appreciated by those skilled in the art that reference herein to treatment extends to prophylaxis as well as the treatment of established conditions or symptoms.

It will be further appreciated that the amount of INFx inhibitor of the invention required for use in treatment will vary not only with the route of administration but also with the nature of the condition being treated and the age and condition of the patient and will be ultimately at the discretion of the attendant physician or veterinarian. In general however, a suitable dose will be in the range of from about 5.0 to 500 up per kilogram of bodyweight per day,

for example, in the range 30 to 300 μ g/kg/day, preferably in the range 50 to 150 μ g/kg/day.

The desired dose may conveniently be presented in a single dose or as divided doses administered at appropriate intervals, for example, as two, three, four or more sub-doses per day.

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While it may be possible that, for use in therapy, a ${\sf TNF}_2$ inhibitor of the invention may be administered as the raw protein it is preferable to present the active protein as a pharmaceutical formulation.

The invention further provides a pharmaceutical formulation comprising a TNFa inhibitor as herein defined or a pharmaceutically acceptable derivative thereof together with one or more pharmaceutically acceptable carriers thereof and, optionally, other therapeutic and/or prophylactic ingredients. The carrier(s) must be 'acceptable' in the sense of being compatible with the ingredients of the formulation and not deleterious to the recipient thereof.

The inhibitors according to the invention may therefore be formulated for parenteral administration (e.g. by injection, for example bolus injection or continuous infusion) and may be presented in unit dose form in ampoules, pre-filled syringes, small volume infusions or in multi-dose containers with an added preservative. The compositions may take such forms as suspensions or solutions in aqueous vehicles, and may contain formulatory agents such as suspending, stabilising and/or dispersing agents. Alternatively, the active ingredient may be in powder form obtained by aseptic isolation of sterile solid or by lyophilisation from solution, for constitution

with a suitable vehicle, e.g. sterile, pyrogen-free water, bofore use.

The TAF: inhibitor of the invention may also be used in combination with other therapeutic agents, for example, other cytokines or inhibitors thereof.

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The invention thus provides, in a further aspect, a combination comprising a TNFx inhibitor as herein defined or a pharmaceutically acceptable derivative thereof together with another therapeutically active agent, for example, other cytokines or inhibitors thereof.

The proteins of the invention may be prepared by purification from natural sources and, where appropriate followed by chemical modification, or they may be prepared by conventional methods known in the art for the preparation of proteins, for example, by recombinant DNA techniques.

According to a further aspect of the present invention, there is provided a process for producing the tumour necrosis factor π inhibitor of the invention by purification from natural sources, particularly the unine of human febrile patients. Such purification, for example, comprises the steps of concentrating the crude unine of febrile human patients, precipitating crude INFα INH from the unine and fractionating the TNFα INH from the other proteins of this precipitate by one or more of, for example, ion exchange column chromatography, gel filtration chromatography, hydrophobicity chromatography, immunoabsorption and affinity chromatography on immobilized INFα.

The tumour necrosis factor a inhibitor of the invention is also obtainable from macrophage containing human tissue, for example,

lung lavages and extracts of human liver, from which it may be obtained by standard purification techniques such as those described above.

Natural and recombinant TNFa INH produced according to the procedures described herein may be purified through a series of steps as listed above. After each of the purification steps, the presence and purity of the INFo INH may be measured in an assay of cytotoxicity in the presence of actinomycin D (acti D) using a TNF-susceptible cell line L929, as described by G. Nedwin et al., J. Immunol., 135, loc. 10 cit.

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In a preferred embodiment of the process the TNF α IN β is initially isolated from untreated urine collected from febrile human patients (>38.5°C) devoid of urinary infections using a standard concentration technique, for example, ultrafiltration. A crude fraction may then be precipitated from the crude urine using ammonium sulphate e.g. by addition of ammonium sulphate to a concentration of 80% (w/v) at 4^{0} C with stirring. Preferably the ammonium sulphate max be added in a stepwise manner and material precipitated at lower concentrations e.g. at 40% (w/v) discarded. The ammonium sulphate may 20 be removed by dialysis and the resulting fraction purified to secarate the TNFa INH from other proteins by a variety of chromatographic methods.

Thus, the TNF INH concentrate may be purified by ion-exchange chromatography which separates proteins according to their differences 25 in electrical charge, which is a reflection of the acid/base properties of the proteins. Suitable materials for anion-exchange chromatography include aminoethylcellulose derivatives, for example,

quaternary-aminosthylcellulose (QAE-cellulose) or diethylaminosthyl-cellulose (DEAE-cellulose) which are widely commercially available.

The anion-exchange column should be equilibrated prior to ecolying the concentrate using a suitable buffer, such as Tris-HCl, optionally containing a chelating agent such as EDTA. Bound material may be eluted from the column using a salt solution (for example, 0.8M sodium chloride made up with the equilibration buffer).

The pooled active fractions from the anion—exchange chromatography chromatography. Suitable materials for cation—exchange chromatography include derivatives of cellulose such as carboxymethyl (CM) cellulose or Sulphopropyl Sepharose (Pharmacia, Uppsala, Sweden). The column should be equilibrated with a suitable buffer, such as sodium acetate and bound material may be eluted with the equilibration buffer containing, for example, 0.5M sodium chloride.

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The pooled active fractions are further purified by affinity chromatography on bound recombinant human TNFα (rhTNFα), coupled to a suitable matrix, for example, Mini-Leak Agarose (Kem En Tec, Biotechnology Corp., Denmark). The column should be buffered using, for instance, a phosphate buffer (e.g. 0.8M potassium phosphate pt 20 8.6). Active groups not bound to rhTNFα should be blocked using ethanolamine-HCl pH 8.5 buffer. The column should be equilibrated with a suitable buffer, for example, Tris-HCl optionally containing sodium chloride and the TNFα INH eluted with an acidic (pH 3.5) glycine buffer. The eluted fractions should immediately be balanced to pH 7.0 by the addition of, for example, Tris base.

The active pooled fractions are preferably lyophilised prior to the final purification step of reverse-phase FPLC (fast protein liquid

chromatography). Prior to its application to the FPLC column, the lyophilised TNFx INM fraction should be buffered with an appropriate buffer such as trifluoroacetic acid (TFA) (Fulka, Buchs, Switzerland), heptafluorobutyric acid (HFBA) or acetic acid. Elution of the TNFx INH from the FPLC column may be carried out using conventional techniques, for example, made up with a suitable buffer as previously described, optionally containing an alcohol, Thir is stance, N-propagol.

The eluted fraction should immediately be buffered with, for example, ammonium bicarbonate and lyophilised. The INFT INH is now in a substantially homogenous form suitable for further assaying of biological activity and for preparation in an appropriate form for therapeutic use.

There is thus provided as a further or alternative aspect of the invention a protein which selectively inhibits TNFα-mediated activity which is substantially identical to that obtained in the above process.

The ability to purify TNFz INH of the present invention to homogeneity has enabled the sequencing of the N-terminal portion of this protein molecule. This sequence will assist in the cloning of a gene coding for the TNFa INH thus enabling the production of large quantities of TNFa INH in pure form for further biological investigation and eventually for therapeutic testing and use.

Samples of the homogeneous TNF2 INH of the present invention can

be sequenced by conventional techniques, for example, using a

commercially available automated sequencer employing either ninhydrin
or gas phase detection. The first 17 residues of the amino terminal

portion of the human $\mathsf{INF}\alpha$ INH as herein defined comprises the following sequence :

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Asp-Ser-Val-Cys-Pro-Gln-Gly-Lys-Tyr-Ile-His-Pro-Gln-Cys-Asn-Ser-Ile (identified using an automated sequencer, Model 4774 from Applied Biosystems).

It is further believed that the next three amino acids provide a glycosylation site and that the sequence thus continues

Asn-Ser-Thr-Lys.

It will be appreciated that one potential method of providing large quantities of INFx INH in pure form is through recombinant DNA techniques which are well known in the art. However, the successful use of such techniques requires not only that the natural and recombinant INFx INH or its activity be accurately measured, but also that both the natural and recombinant product be purifiable to homogeneity.

According to yet another aspect of the present invention, there is provided a process for producing a tumour necrosis factor inhibitor of the invention or a derivative thereof by the expression of a DNA sequence encoding for such an inhibitor in a suitable transformed host. Such a process involves the culturing of a host transformed with recombinant DNA molecules comprising DNA sequences encoding the inhibitor which have been inserted into a suitable vector.

Suitable eukaryotic and prokaryotic hosts may be, for example, strains of bacteria, yeasts, other fungi, and animal cells (including insect cells) and plant cells in tissue. Particularly preferred host

cells are yeast cells, E. coli cells and animal cells.

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Expression of a protein having tunour necrosis factor inhibitor activity is achieved by culture of the transformed host cells in a suitable growth medium. Normally such a medium will contain a source of nitrogen such as ammonium sulphate, a source of carbon and energy such as glucose or glycerol, trace elements and factors essential to growth of the particular host cells. The precise culture conditions will be dependent upon the chosen host; thus, for example, in the case of \underline{E} , coli submerged aerobic fermentation is preferred, preferably at about $37^{\circ}C$.

In addition, expression may be induced, for example, by the addition of an inducer or the use of inducing conditions for the promoter system being used in the expression vector.

Depending upon the host, the TNFx inhibitor may be produced as granular inclusion bodies which can be recovered, after cell lysis, by differential centrifugation; these can be solubilised by conventional methods and purified by the methods described herein for purification of urinary TNFα INH. Alternatively, the TNFα inhibitor may be in solution in the cytosol, secreted into the periplasmic space or conveniently secreted into the culture medium.

The host cells are transformed by recombinant DNA molecules which comprise a DNA sequence encoding for a $TNF\alpha$ inhibitor which has been inserted into an expression vector.

Such expression vectors may consist of segments of

chromosomal, non-chromosomal and synthetic DNA sequences, such as

various known derivatives of SV-40 and known bacterial plasmids, for
example, "natural" plasmids such as ColEl, pSCIOI or pRSF2124 and

phage DNAs, or "artificial" plasmids (constructed <u>in vitro</u>) such as p8R322, pM89 or p4T153. Phage DN4s include, for example, the numerical derivatives of phage lambda and other DNA phages, for example M13, and other filamentous single-stranded DNA phages. Vectors useful in yeasts include the 2r plasmid, and those useful in eukaryotic cells such as animal cells include those containing SV-40 adenovirus and retrovirus.

Such expression vectors may also be characterised by at least one expression control sequence which may be operatively linked to the TNF inhibitor DNA sequence such that it controls and regulates the expression of the cloned DNA sequence. Examples of useful expression control sequences include the <u>lac</u>, <u>trp</u>, <u>tac</u> and <u>trc</u> systems, major operator and promoter regions of phage \(\lambda\) (such as the P_L cromoter under the control of the thermolabile <u>ts</u> cl857 repressor), the control region of fd coat protein, the glycolytic promoters of yeast (e.g. the promoter for 3-phosphoglycerate kinase), the promoters of yeast acid phosphatase (e.g. Pho 5), the promoters of yeast \(\alpha\)-mating factors, and promoters derived from polyoma, adenovirus, retrovirus, and simian virus.

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In addition, such expression vectors may possess various sites for insertion of the TNF α inhibitor DNA sequences of this invention. These sites are characterised by the specific restriction endonuclease which cleaves them. Such cleavage sites are well recognised by those skilled in the art. The expression vector, and in particular the site chosen therein for insertion of a selected DNA fragment and its operative linking to an expression control sequence, is determined by a variety of factors including the number of sites susceptible to a

given restriction enzyme, the size of the protein to be expressed, contamination or binding of the protein to be expressed by most definition which may be difficult to remove during purification, the location of start/stop codons, and other factors recognised by those skilled in the art. Thus the choice of a vector and insertion site for a DNA sequence is determined by a balance of these factors, not all selections being equally effective for a given case.

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Likewise, not all host/vector combinations will function with equal efficiency in expressing the DNA sequences of this invention. The selection is made, depending upon a variety of factors including compatability of the host and vector, ease of recovery of the desired protein, expression characteristics of the DNA sequences and the expression control sequences operatively linked to them, or any necessary post-expression modifications of the desired protein.

The DNA sequences of the invention which on expression code for proteins with TNFa inhibitor activity may be isolated by screening various DNA libraries for such DNA sequences using a series of DNA probes. The DNA probes may be prepared from the purified natural protein which is used as a source of amino acid sequence data. The purified natural protein may be prepared, for example, from febrile human urine as described above. Degenerate DNA sequences coding for various portions and fragments of the amino acid sequence, e.g. in combination with Lathe probes, are used to design the DNA probes.

Thus, various DNA libraries are screened for DNA sequences coding for the INFa inhibitors of the invention. Such libraries include chromosomal gene banks and cDNA or DNA libraries prepared from cell lines or tissue that are demonstrated to produce INFa

inhibitors, such as alveolar macrophages or liver tissue. Screening may be by direct immune expression, for example in igill or similar systems, or, in the case where a TVFx INH producing cell is identified, by identification of TVFx INH specific mRNA by direct expression in Xenopus pocytes.

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A variety of conventional cloning and selection techniques may be used to locate and identify DNA sequences (M.C.) encode on expression in an appropriate eukaryotic or prokaryotic host for the TNFa inhibitors of this invention. These selected DNA sequences may themselves be used as probes to select other DNA sequences coding for TNFx inhibitors or may be used in appropriate recombinant DNA molecules to transform appropriate eukaryotic or prokearyotic hosts for the production of TNFx INH encoded by them.

The invention includes within its scope single and double stranded DNA sequences encoding for TNF2 INHs, vectors containing such sequences suitable for transformation of a host organism and host cells transformed with such DNA sequences.

According to a further aspect of the present invention we provide a protein with selective TNFx inhibitor activity produced by expression of a host transformed with a DNA sequence encoding for such a TNFx inhibitor protein. TNF inhibitors of the invention which are prepared by the expression of a DNA sequence encoding such inhibitors in a transformed host will thus be identical to the sequence of native TNFx INH or contain one or more deletions, substitutions, insertions, inversions or additions of allelic origin or otherwise, the resulting sequence will have at least 90% and preferably 90% homology with the sequence of native INFx INH and retain essentially the same biological

properties. In particular, a INF inhibitor of the invention may include an N-terminal methionine. Also, for example, the DNA sequence of the invention coding for INFa INH may be fused in an expression vector to a portion of a DNA sequence coding for a eukaryotic or prokaryotic polypeptide to assist the expression of the INFa INH encoding DNA sequence or aid secretion, maturation or purification of the INFa INH from the host; the fused polypeptide may be removed intra-or extra-cellularly by known techniques or the INFa INH may be used together with the fused polypeptide.

The INF α INHs produced by culturing of the eukaryotic and prokaryotic hosts transformed with DNA sequences encoding for INF α INHs can then be employed, after purification, in the pharmaceutical compositions of this invention.

It will be appreciated that, when produced by animal cells, the INF INF INF INF invention will be a glycoprotein. Prokaryotic expression systems will, however, produce the protein in an unglycosylated state. In addition, the glycosylated protein may be substantially deglycosylated by techniques known in the art, for example, by the use of endoglycosidase enzymes.

The following non-limiting Examples illustrate the invention. All temperatures are in ${}^{\circ}$ C and all percentage concentrations are w'.

TNFc Inhibition Assav

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The percentage of TNFα INH activity in the fractions described in the Examples was determined by assuming that the optical density (OD) values from murine L929 cells stimulated by actinomycin D (actinophycin D) corresponded to 100% inhibition, whereas the OD from cells cultured

with actinomycin D and INFx corresponded to a maximal cell mortality of 0% INFx inhibition. The INFx used in the assay was recombinant human INFx (rhINFx) produced in <u>E. coli</u> as described by 4. Marrenout etc. al., "Molecular cloning and expression of human tumbur necrosis factor and comparison with mouse tumbur necrosis factor", <u>Eur. J. Biochem.</u>, 152, p515 (1985). Thus the percentage of INFx inhibition in the assay of cytotoxicity was calculated according to formula (I)

Percentage of INFx INH activity =

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Example 1

Purification of Urinary TNF2 INH

a) Concentration of Protein from Human Urine

five patients prior to any treatment. Two of the patients were suffering from small-cell carcinoma, one from malignant histocytosis, one from polymyocitis and one from sepsis. All were highly febrile (>38.5°C) and devoid of uninary infections. The unine was concentrated at 40 on an Amicon ultrafiltration hollow fibre apparatus, with a molecular size cut-off of about 5 kDa.

b) Precipitation of Protein from Human Urine

The concentrated unine pool was saturated with solid graphics sulphate by adding the sulphate slowly with constant stirring at 4-until an ammonium sulphate concentration of 40% was reached. The precipitate was removed by centrifugation, discarded, and the supernatant adjusted to 80% saturation with addition of further ammonium sulphate. A pellet was obtained by centrifugation which was resuspended in 150 ml of 20 mM additum phosphate (pH 7.2) and 150 mM sodium chloride. The ammonium allphate was removed by dialysis at 4-using 10 mM Tris-HCl pH 7.4, 2 mM EDTA and 5 mM benzamidine FCL.

c) Identification of TNFx INH Activity

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The semi-purified fraction of Example 1(b) was tested in a cytoxicity assay with the INF-susceptible cell-line L929 in the presence of actinomycin D. At a 1:20 dilution of the semi-purified fraction, total inhibition of the cytotoxic effect induced by $rhTNF_2$ was observed so that the $OD_{570\,\mathrm{nm}}$ value was identical to that measured in the presence of actinomycin D alone (OD $_{570\,\mathrm{nm}}=1.5$).

Furthermore, inhibitory activity was observed in dilutions of the fraction of up to 1:160 on cells $(00_{370}\,\mathrm{nm}=0.83)$ whereas the control value of rhTNF α at a final concentration of 0.2 ng/ml measured in the presence of actinomycin D was lower $(00_{370\,\mathrm{nm}}=0.73)$, so that 50% of inhibition was observed at a dilution of approximately 1:190 $(00_{370\,\mathrm{nm}}=1.10)$. The TNF α INH had no effect on cell viability when tested without actinomycin D.

d) Compension of the Effect of TWE 1NH on TWE 1 and 3 Induced Cytotoxicity

A cytotoxicity assay was conducted using the INF-susceptible cell-line L929 in the presence of actinomycin D using a range of concentration of INFx or INF3 to induce the cytotoxic effect. The semi-purified fraction from Example 1(b) was tested at 1:20, 1:50 and 1:90 dilutions. Control tests were performed in the absence of the INFx or INF8 cytokine, and in the absence of inhibitor. The results are shown in Table 1, below, and demonstrate that the inhibitor of the invention does have some inhibitory effect on INF3 mediated cytoxicity ranging from approximately 50% down to 2% of the INFx inhibition with increasing INF5 concentration.

TABLE 1

Final Concentration of TNF (x or 3) (pg/ml) Added to Actinomycin-D Treated L929 cells	Form of Cytokine acded to cells	Final Dilution of Secherryl S-200 Innibitory Fraction or L929 cells (OC ₁₇₁ nm)			
		None	1/20	1 ′53	1/50
0	0	>1.90	>1.90	>1.90	>1.90
10	a	V.O	NO _c .	ND	NC.
	β	1.16	1.66	1.46	1.39
20	α .	1.30	>1.90	1.72	1.66
	3	1.02	1.51	1.21	1.22
50	Œ	1.02	>1.90	1.72	1.é9
	,s	0.65	1.03	0.84	0.55
100	Œ	0.73	1.78	1.69	1.51
	3	0.37	0.71	0.59	0.52
250	Œ	0.38	1.70	1.70	1.33
	3	0.24	0.44	0.31	0.24
500	Œ	0.30	1.52	1.49	1.11
	9	0.17	0.30	0.26	0.18
1 250	· α	0.19	1.09	1.10	0.7é
	۶	0.11	0.13	0.19	0.13
2 500	α	0.06	1.03	0.80	0.47
	9	СИ	ND	פא	ND

Example 2

Gel Filtration of Uninery INF: INH

The semi-purified TNFx INH of Example 1(b) was purified by gel filtration chromatography at 4° on a Sephachyl S-200 column (0.9 x 60 cm) (Pharmacia, Uppsala, Sweden) equilibrated in 50 mM Tris-HCL buffer (pH7.4) containing 100 mM sodium chloride. A sample of the protein fraction (20 mg, 0.8 ml) was applied to the column and eluted with the same buffer at a flow-rate of 5.4 ml/hr. Fractions (1.35 ml) were collected and tested for TNFx INH activity. The TNFx INH activity eluted from the gel in a single peak. The inhibitory activity showed an apparent molecular weight of 40 to 60 kDa (see Figure 1).

Example 3

Chromatofocussing of Urinary TNF: INH

The semi-purified TNFx INH of Example 1(b) was chromatofocussed at 4° on a Mono-P pre-packed column (HR 5/20, 5 x 200 mm) (Pharmacia, Uppsala, Sweden) equilibrated in 25 mM Bis-Tris buffer adjusted to pH 7.1 with imidodiacetic acid (Fluka, Buchs, Switzerland). A sample of the protein fraction of Example 1(b) (30 mg) was applied to the column and eluted with a polybuffer 74/iminodiacetic acid at pH 4.0. Column fractions (1 ml) were tested at 1:10 dilution for their effect in the rhTNFa (0.2 ng/ml) cytotoxicity assay in the presence of actinomycin D (lug/ml). The actual pH of each column fraction was determined with a pH meter, the bulk of the TNFa INH activity being contained in the eluted fractions

between pH 5.5 and 6.1 (see Figure 2). This is equivalent to the pI of the TNFs INH protein.

Example 4

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5 Ion-Exchange Chromatography of Uninary INF: INH

The semi-purified INFz INH of Example 1(b) was purified by anion-exchange chromatography at 4° on a CEAE Sephadex column (2.6 x 20cm) (Pharmacia, Uppsala, Sweden) equilibrated in 10mM Tris-HCl buffer pH 8.0, containing 2mM EDTA. Bound material was eluted from the column with the equilibration buffer containing 0.8M sodium chloride. Fractions (8.0ml) were collected, tested for INFz INH activity and the inhibitory fractions were pooled (160ml) and dialysed against 10mM sodium acetate buffer pH 5.0 (4 x 2 litres).

The TNFα INH was further purified by cation—exchange

chromatography at 4° on a Sulphopropyl—Sephadex column (0.8 x 15 cm)

(Pharmacia, Uppsala, Sweden) equilibrated in 10mM sodium acetate

buffer pH 5.0. Bound material was eluted from the column with the

equilibration buffer containing 0.5M sodium chloride. Fractions

(7.5ml) were collected, tested for TNFα INH activity and the

inhibitory fractions were pooled and concentrated 20-fold on an Amicon

ultrafiltration apparatus with a molecular size cut-off of about 10

kDa.

Example 5

25 Gel Filtration of Uninary INF: INH

The TNF α INH concentrate from Example 4 was purified by gel filtration chromatography at 4 $^{\circ}$ on a Sephacryl S-200 column (2.5 x 100

cm) (Pharmacia, Uppsala, Sweden) equilibrated with SOme Tris-ICL pm 7.4 buffer containing 100mH sodium chloride. A sample of the protein fraction from Example 4 (200mg) was applied to the column and eluted with the equilibration buffer at a flow-rate of 27m1/hour. Fractions (9.0ml) were collected, tested for INFx INH activity and the inhibitory fractions were pooled. The column was calibrated with dextran blue (DB), 2000 kDa; bovine serum albumin (BSA), 67 kDa; ovalbumin (OA), 43 kDa; α -chymotrypsinogen-A (α CT), 25 kDa; and ribonuclease A (RNase), 13.5 kDa, as shown in Figure 4.

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Example 6

Affinity Chromatography of Urinary TNFx INH

A TNFx affinity column was prepared by coupling recombinant human INFx (1.0mg) to Mini Leak Agarose (Kem En Tec, Biotechnology Corp.,

Denmark) in 0.8M potassium phosphate buffer pH 8.6. The remaining active groups were blocked by incubation in 0.1M ethanolamine-HCl buffer pH 8.5. The gel was washed with 50mM Tris-HCl pH 7.4 buffer containing 100mM sodium chloride (3 x 50ml). A sample of the TNFx INH fractions from Example 5 (15ml) was applied to the column and eluted with a 0.2M glycine-HCl pH 3.5 buffer. Fractions (1.0ml) were collected, immediately adjusted to pH 7.0 by addition of 1M Tris (5 to 40ml) and tested for TNFx INH activity.

Example 7

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Reverse Phase FPLC Chromatography of Urinary TNF & INH

The TNFa INH fractions from Example 6 were lyophilised, dissolved in 0.1% trifluoroacetic acid (2.0ml) and loaded onto a ProRPC

reverse-phase FPLC column (5 x 20 cm) (Pharmacia, Uppsala, Swegen) equilibrated in 0.1% trifluoroacetic acid. Bound material was eluted with a 0 to 100% acetonitrile gradient in 0.1% trifluoroacetic acid at a flow rate of 0.3ml/minute. To each fraction (0.75ml) 0.5M ammonium bicarbonate (10 μ 1) was added and the eluted material was lyophilised.

The reverse-phase FPLC chromatography revealed one major peak corresponding to TNFz INH activity. The lyophilised fractions containing this activity were dissolved in 10mM Tris-HCl pH 7.4 buffer containing 2mM EDTA and analysed by sodium dodecyl sulphate polyacrylamide gel electrophoresis (SDS PAGE) using the method described by U. Laemmli et al., Nature, 277, p 680 (1970). The TNFa INH was found to elute with a molecular weight of 33 kDa (see Figure 4). Samples run under non-reducing conditions were tested for TNFz INH activity at 1:10 dilution on L929 cells in the presence of 0.15 mg/ml of rhTNFa. The activity directed against rhTNFz migrated with an apparent molecular weight identical to the 33 kDa band on the gel run under reducing conditions.

Example 8

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Protein Sequencing of Urinary TNF INH

The TNF¢ INH fraction isolated from the reverse-phase FPLC chromatography was concentrated in vacuo and spotted onto a conditioned sequencer filter. The protein was analysed with an Applied Biosystems Model 477A protein sequencer. Fractions from the sequencing cycles were evaporated to dryness and resuspended in N,N-disopropylethylamine acetate and acetonitrile prior to injection into an HPLC column for residue identification.

The first 17 amino acid residues of the N-terminal were identified and have the sequence: Asp-Ser-Val-Cys-Pro-Gln-Gly-Lys-Tyr-Ile-His-Pro-Gln-Cys-Asn-Ser-Ile. It is further believed that the next three amino acids provide a glycosylation site and that the sequence thus continues Asn-Ser-Thr-Lys. This sequence is not significantly homogeneous to any protein sequence contained in the NBRF Protein Sequence database (November 1988).

Example 9

10 Demonstration that INF: INH is a Protein

a) Time and Temperature Dependency

The Sephacryl S-200 purified INFx INH of Example 2, obtained by bulking the tubes of the active fractions, was heated at 56°, 75° and 95°. The INFx INH activity was measured after 10, 20 and 60 minutes and, by comparison with untreated samples, the percentage of INFx INH activity was calculated according to formula (I). The results shown in Table 2 below demonstrate that the INFx INH activity decreases in a time- and temperature-dependent manner.

Heat inactivation					
Temperature (°C)	Time (min)	Percentage of INFo INH activity			
56	10 20 60	100 100 93			
75	10 20 · · · · 60	60 26 15			
95	10 20 60	27 10 13			

b) Susceptibility to Trypsin Digestion

Trypsin (500µg) (Sigma, St. Louis, MO) in 0.2M Tris-HC1 buffer (pH 8.0) containing 1mM calcium chloride was added to the pooled fractions of Sephacryl S-200 purified urinary TNF α INH of Example 2 and incubated at 37°C for 4 hours. Another measure of trypsin (500 α) was added and digestion continued for a further 20 hours, at which time the reaction was terminated by addition of soybean trypsin inhibitor (2mg) (Sigma, St. Louis, MO). The percentage of INF α INH inhibitory activity of the trypsin digest and the control was determined at a 1:20 final dilution of the pool of bulked fractions on L929 cells stimulated by rhINF α in the presence of actinomycin D, according to formula (I). The results are shown in Table 3 below.

Trypsin inactivation					
Conditions	Pecentage of TNFx I\H activity	· CD 570nm			
Buffer alone	0	0.71			
Trypsin + soybean trypsin inhibitor in buffer	o • ·.	0.70			
Partially purified Sephadex S-200 urine	61	1.46			
Partially purified Sephadex S-200 urine digested by trypsin	23	1.03			

c) Treatment with Urea

The Sephacryl S-200 purified TNF α INH of Example 2 was adjusted to 2M urea and extensively dialysed at ϵ^{μ} against phosphate buffered saline (PSS) containing 2M urea. Dialysis against PES was repeated prior to the bioassay. TNF α INH activity was found to be unaffected which indicates that inhibitory activity is not mediated by a molecular weight bound to a protein.

Example 10

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Demonstration of Competitive Inhibition

The Sephacryl S-200 purified TNFa INH of Example 2 was tested at a 1:10 dilution against increasing amounts of rhINFa on L929 cells. An inverse correlation between the amount of rhINFa present in the assay and the degree of inhibition was observed (see Figure 3). Thus, the inhibitory activity is competitively overcome by increasing concentrations of rhINFa.

Example 11

Inhibition of INFx-Mediated PGE, Production by Dermal Fibroblasts

Human dermal fibroblasts were seeded at a concentration of 2.0 x 10° cells/well and cultured for 48 hours. Cells were then stimulated with DMEM buffer supplemented with 10% FCS as a control. Cells were also stimulated with rhINF at concentrations ranging from 0.5 to 5mg/ml, and the effect of the TNF α INH from Example 5 was studied at three dilutions (1:20, 1:50 and 1:80) in the above buffer. After 72 hours of incubation, PGE2 production was measured in the supernatants by radioimmunoassay using a PGE2 antiserum [see J M Dayer et al., J. Clin. Invest., 67, p1386 (1979)].

The results are shown in Table 4, below, and demonstrate that the ability of rhTNF α to stimulate PGE $_2$ production by dermal fibroblests was inhibited by the addition of TNF α INH at all three dilutions. At 1:80 dilution of TNF α INH the inhibitory activity was partially overcome by increasing rhTNF α concentrations.

Concentration of rhTNF on human fibroblasts	PGE ₂ production by human dermal fibroblasts (ng/ml) Dilution of TNF INH on fibroblasts				
(pg/ml)	none	1:80	1:50	1.11	
0	50.6 ± 7.4	86.8 • 5.6	103.0 • 8.9	111.0 - 9.4	
500	160.0 <u>•</u> 14.1	126.0 - 9.3	115.9 + 6.6	113.9 - 7.1	
2,000	331.7 <u>+</u> 28.4	217.2 <u>+</u> 10.7	156.7 ± 10.7	115.3 ± 21.3	
5.00G	381.7 ± 19.6	257.2 ± 13.7	253.1 <u>•</u> 21.2	221.6 <u>•</u> 16.0	

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Three different experiments were carried out with the same strain of fibroblasts. Guffer or TNFG INH was incubated at various dilutions in the presence or absence of various concentrations of rhTNFG. PGE2 production by cultured human dermal fibroblasts was neasured after three days. Values represent triplicate means of the three cultures • SEM (N=3).

Example 12

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rhTNFa Binding Inhibition Assay

Recombinant human TNF α was indinated by using the indocen method of Fraker and Speck jr., Biochem. Biophys. Res. Comm., 80, p849 (1978). The specific activity of [125 I]-TNF α was 2.2x10 4 cpm/ng and produced a single band with a molecular weight of 17 kDa when analysed by SDS PAGE. The human macrophage cell line U937 in aliquots of 10^6 cells was cultured at 4^{0} for 2 hours in a culture medium (200 μ 1) comprising RPMI 1640 (Gibco, Paisley, Scotland) supplemented with streptomycin (100 μ g/ml), penicillin (100U/ml), 1.0% glutamine and 10% foetal calf serum, and additionally containing 0.04% sodium azide and 0.5ng [12 I]-TNFa. Binding inhibition was performed by the addition of various dilutions of TNE INH $(1:20,\ 1:200\ and\ 1:2000)$.

Non-specific binding was measured in the presence of a 100-fold excess of unlabelled $rhTNF\alpha$, and free readioactivity was separated from the bound $[^{1.2i}$ I]-TNFlpha by centrifugation through an bil mixture as described by Robb et al., J. Exp. Med., 154, p1455 (1981). Cell bound [12 I]-TNR was measured in a gamma counter (LKB, Bromma, Sweden), and the percentage of binding inhibition was determined according to formula (II)

25 100 x 1 - cpm with TNF INH - cpm non-specific binding cpm of total binding - cpm non-specific binding (II)

Percentage of binding inhibition =

Example 13

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Effect of TNE INH on [12 I]-TNE2 Binding to U937 Cells

U937 cells were preincubated for 1 hour at 20° in the culture medium of Example 12, in the presence of either [12 I]-TNF2 alone or [125 I]-TNF2 with a 100-fold excess of unlabelled rhINF2. The U937 cells were then washed with phosphate buffered saline (3 x 50ml) at 40 . The cells incubated in [125 I]-TNF2 alone were divided into four batches and incubated with TNF2 INH from Example 5 (1:20, 1:200 and 1:2000 dilutions) and with buffer alone, respectively.

The specific binding of [125 I]-INFx to U937 cells was found to be inhibited at 40 by 100%, 80% and 35% by the three dilutions of INFx INH, 1:20, 1:200 and 1:2000, respectively (see Figure 5). The control batch which lacked INFx INH showed no inhibitory activity. The binding inhibition of the two weaker dilutions was found to be increased to 90% and 60% when [125 I]-INFx was preincubated with INFx INH at dilutions 1:200 and 1:2000, respectively, prior to cell addition.

The experiment was repeated using the cells preincubated in the presence of $[^{12}$ I]-TNF α and a 100-fold excess of unlabelled TNF α so that the percentage of binding inhibition could be corrected for non-specific binding.

Example 14

Dissociation of a Pre-Formed TNFa: U937 Complex

U937 cells were preincubated for 1 hour in the presence of $[^{125}\,{}^{\prime}I]$ -TNF α as described in Example 12. The cells were washed and incubated at either 4^0 or 37^0 in the presence or absence of TNF α INH

from Example 5. Cell-surface bound [-251]-TAFa was found to dissociate faster in the presence of TAFa INH than in its absence and this was found to occur in a time- and temperature-dependent manner (see Figure 6).

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Example 15

Demonstration that INF2 INH is not Protective for the INF2

[12 I]-INFx was incubated at 20 for 1 hour in the presence of 3 different dilutions of INFx INH (1:20, 1:200 and 1:2000) and in the presence of buffer alone. When analysed by SDS PAGE and autoradiography the [125 I]-INFx was found to migrate as a single band both in the absence and presence of INFx INH showing the inhibitor to have no protectytic effect.

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Example 16

Effect of TNFs INH on IL-1 Receptor Binding Activity

The activity of TNFa INH from Example 5 was tested in the IL-1/LAF (lymphocyte activating factor) assay when induced by IL-12 or IL-13 [this assay is described by P. Seckinger et al., J. Immunct., 139, p1541 (1987) for an IL-1 inhibitor protein]. A dose response of [3H]-thymidine incorporation (corresponding to thymocyte proliferation) was observed in up to 200pg/ml concentrations of both IL-la and IL-13. Addition of TNF INH at levels observed to inhibit rhINFa did not have any significant effect on the IL-1-induced thymocyte proliferation, proving inhibition to be specific for INF2 only.

The results obtained are illustrated by reference to the accompanying drawings, in which:-

Figure 1 - shows the uninary TNFα INH activity profile of

Sephacryl S-200 gel filtration. Column fractions (lml) were tested at

1:10 dilution for effect in the rhTNFα (l.Ong/ml) cytotoxicity assay
in the presence of actinomycin D (l.Oμg/ml) (o—o). The line

(—) represents OD_{280 nm} of the fractions. Bars represent cell

lysis measured by dye uptake in response to actinomycin D (□) and to

actinomycin D plus hrTNFα (□) without unine. The molecular weight
markers are dextran blue (D8), bovine serum albumin (BSA), ovalbumin

(OA), α-chymotrypsinogen (αCT), ribonuclease A (RNase) and phenol red

(O-red).

Figure 2 - shows the uninary TNFα INH activity profile of chromatofocussing on a Mono-P column. Column fractions (lml) were tested at 1:10 dilution for effect in the chTNFα (0.2ng/ml) cytotoxicity assay in the presence of actinomycin D (1.0 μg/ml) (----). The line (----) represents their pH. The bars are as described for Figure 1.

Figure 3 - shows the reversibility of TNF α INH activity. Open circles (o—o) represent $OD_{570\,\text{nm}}$ measured in the presence of actinomycin D, rhTNF α with TNF α INH; closed circles (•—•) represent $OD_{70\,\text{nm}}$ measured in the presence of actinomycin and rhTNF α only and the bar (\square) represents $OD_{570\,\text{nm}}$ in the presence of actinomycin D (1.0 μ g/ml) alone.

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Figure 4 - shows the elution profile of Sepacryl S-200 gel filtration with purified INF α INH from Example 5. Column fractions (9ml) were sterilized and tested at 1:50 dilution against rhINF α (1.0mg/ml) in the presence of actinomycin 0 (1.0ug/ml) in the presence of actinomycin 0 (1.0ug/ml) in the L929 cytotoxicity assay (α). The line (_____) represents 00_{280} nm of the fractions. The bars are as defined for Figure 1.

Figure 5 - shows the SDS PAGE analysis of purified TNFz INH of Example 7. SCS PAGE was performed as described by U. Laemmli et al., Nature, 277, loc cit. Samples were loaded onto 15% polyacrylamide gel with a 3% stacking gel and gels were silver-stained as described by C. Merril et al., Proc. Natl. Acad. Sci. USA, 76, p4335 (1979). Samples run under non-reducing conditions were tested for biological activity by cutting 2mm slices from the gel and eluting the proteins by overnight incubation in 10mM Tris-HCl pH 7.4 containing 2mM EDTA (total volume 200μ1). Fractions were tested at 1:10 dilution on L929 cells in the presence of rhTNFα (0.15ng/ml).

Figure 7 - shows the dissociation of a pre-formed INFz: U937 complex. U937 cells were pre-incubated with [123]-INFx, washed and incubated with INFx INH as described in Example 14, at either 40 ([123]) or 370 ([123]). At the time indicated, cell associated radioactivity was measured and percentage specific binding determined. On the graph, the value obtained from the control without the inhibitor has been substracted from the values obtained at the two temperatures, thus, 100% corresponds to the value obtained without the addition of INFx INH.

CLAIMS

- 1. A protein which inhibits tumour necrosis factor (TNF) α -mediated activity but does not block other proteins which have in common with TNF certain but not all of the biological activities of TNF.
- 2. A protein which selectively inhibits tumour necrosis factor α -mediated activity and having one or more of the following characteristics :
- (a) a molecular weight in the range 40 to 60 kDa, determined by molecular sieve chromatography;
- (b) an iso-electric point (pI) in the range 5.5 to 6.1, determined by chromatofocussing;
- (c) inhibition of the standard TNF assay of differential cytotoxicity for murine L929 cells treated with actinomycin D;
- (d) inhibition of TNF-induced PGE_2 release from human fibroblasts and synovial cells;
- (e) the inhibitor interferes with the binding of $TNF\alpha$ to U937 cells (a monocytic tumour line) as evidenced by inhibition of binding of radiolabelled $TNF\alpha$ (125 I- $TNF\alpha$):
- (f) the dissociation of a pre-formed TNF α : U937 cell complex is promoted by the inhibitor in a temperature dependent manner;
- ____(g) the inhibitor does not degrade TNF by proteolytic. cleavage;
- (h) the inhibitor does not inhibit IL-1 receptor-binding activity e.g. the binding of radiolabelled IL-1 ($^{125}I-IL-1a$) to the murine thymoma subline EL4-6.1.

- 3. A protein which selectively inhibits tumour necrosis factor amediated activity and having one or more of the following characteristics:
- (a) a molecular weight of about 33 kDa, determined by SDS PAGE;
- (b) an iso-electric point (pI) in the range 5.5 to 6.1, determined by chromatofocussing;
- (c) inhibition of the standard TNF assay of differential cytotoxicity for murine L929 cells treated with actinomycin D;
- (d) inhibition of TNF-induced PGE_2 release from Human fibroblasts and synovial cells;
- (e) the inhibitor interferes with the binding of TNFs to U937 cells (a monocytic tumour line) as evidenced by inhibition of binding of radiolabelled TNFs (125 I-TNFs);
- (f) the dissociation of a pre-formed TNF α : U937 cell complex is promoted by the inhibitor in a temperature dependent manner;
- (g) the inhibitor does not degrade TNF by proteolytic cleavage;
- (h) the inhibitor does not inhibit IL-1 receptor-binding activity e.g. the binding of radiolabelled IL-1 (125 I-IL- 12) to the murine thymoma subline EL4-6.1.
- 4. A protein as claimed in either of claims 2 or 3 having the properties (a) and (b) together with one or more of the properties (c) to (h).

- 5. A protein as claimed in either of claims 2 or 3 having all of the properties (a) to (h).
- 6. A protein according to any one of claims 1 to 5 which corresponds to a naturally occurring $TNF\alpha$ inhibitor.
- 7. A protein according to any one of claims 1 to 6 having an amino terminal amino acid sequence as follows:

Asp-Ser-Val-Cys-Pro-Gln-Lys-Tyr-Ile-His-Pro-Gln-Cys-Asn-Ser-Ile.

8. A protein according to claim 7 having an amino terminal amino acid sequence as follows:

Asp-Ser-Val-Cys-Pro-Gln-Gly-Lys-Tyr-IIe-Hts-Pro-Gln-Cys-Asn-Ser-IIe-Asn-Ser-Thr-Lys.

- 9. A protein according to any one of claims 1 to 8 and in which the amino acid sequence contains one or more deletions, substitutions, insertions, inversions or additions of allelic origin or otherwise, the resulting sequence having at least 80% homology with the parent protein and retaining essentially the same biological properties as the parent protein.
- 10. A protein according to claim 9 having at least 90% homology with the parent protein.

- 11. A protein according to any one of claims 1 to 10 in a substantially homogeneous form.
- 12. A protein according to any one of claims 1 to 11 which is a recombinant protein.
- 13. A protein according to any one of claims 1 to 12 which is a glycosylated protein.
- 14. A protein according to any one of claims 1 to 12 which is in a substantially unglycosylated state.
- 15. An exogeneous DNA comprising a nucleotide sequence coding for a protein as defined in any one of claims 1 to 11.
- 16. A cONA comprising a nucleatide sequence coding for a protein according to any one of claims 1 to 11.
- 17. A recombinant expression vector comprising DNA according to either of claims 15 or 16.
- 18. A host cell transformed with an expression vector according to claim 17.

- 19. A method of producing a TNF α INH protein which comprises culturing a cell according to claim 18 and isolating the TNF α INH protein.
- 20. A recombinant protein produced according to the method of claim 19.
- 21. A method for the preparation of a TNF α INH protein comprising the steps of :
 - (a) concentration of urine from febrile patients;
 - (b) ammonium sulphate precipitation;
 - (c) anion-exchange chomatography;
 - (d) cation-exchange chromatography;
 - (e) gel filtration;
 - (f) affinity chromatography; and
 - (g) reverse phase FPLC.
- 22. A protein characterised in that it is substantially identical to the protein obtained by the method of claim 21.
- 23. A pharmaceutical formulation comprising a TNFa inhibitor as defined in any one of claims 1 to 14 or claim 22 or a pharmaceutically acceptable derivative thereof and a pharmaceutically acceptable carrier therefor.

- 24. A protein as defined in any one of claims 1 to 14 or claim 22 for use in therapy.
- 25. A pharmaceutical formulation for use in the manufacture of a medicament for the treatment of conditions associated with excessive or unregulated $INF\alpha$ production, wherein said formulation comprises a $INF\alpha$ inhibitor as defined in any one of claims 1 to 14 or claim 22 or or a pharmaceutically acceptable derivative thereof.